

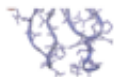
Table

Rank	File ID	Compound	Affinity	Total Energy	vdW Energy	Elec. Energy
1	c359152ed8	<input checked="" type="checkbox"/> ligand 1	-4.830	-86.676	-0.000	0.113
		<input checked="" type="radio"/> run 4	-4.830	-86.676	-0.000	0.113
		<input type="radio"/> run 4	-4.830	-86.675	-0.000	0.114
		<input type="radio"/> run 4	-4.830	-86.675	-0.000	0.114
2	6f9df35049	<input checked="" type="checkbox"/> ligand 1	-4.657	-28.226	-0.000	0.036
		<input type="radio"/> run 12	-4.657	-28.226	-0.000	0.036
		<input type="radio"/> run 12	-4.657	-28.226	-0.000	0.036
		<input type="radio"/> run 12	-4.657	-28.226	-0.000	0.036

3D View

Use your mouse to drag, rotate, and zoom in and out of the structure:

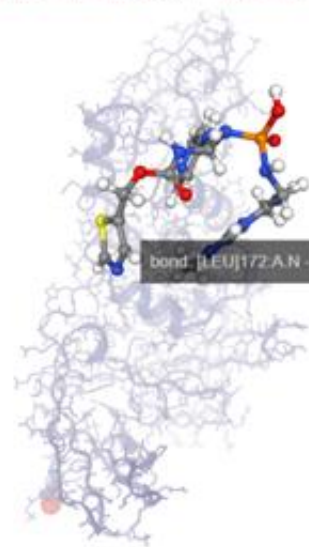
auto view shift + left_mouse + drag zoom ctrl + right_mouse + drag rotate



3D View

Use your mouse to drag, rotate, and zoom in and out of the structure:

auto view shift + left_mouse + drag zoom ctrl + right_mouse + drag rotate



bond: [LEU]172:A:N - [LEU]172:A:CA (protein_b7fb85ceb2_prip.pdb)

.../index.php?tab=DOCKING&page=RESULTS&jobId=6wzu_RocustyrnaTM_gs_convs_1XX2XXpdb_62b306a...

run 12	-4.657	-28.226	-0.000	0.036
run 12	-4.657	-28.226	-0.000	0.036

1

3D View

Use your mouse to drag, rotate, and zoom in and out of the structure:

auto view shift + left_mouse + drag zoom ctrl + right_mouse + drag rotate

